



<110> Bazar J. Fernando

<120> Mammalian Cytokines; Related Reagents and Methods

<130> DX0903K

<140>

<141>

<160> 9

<170> PatentIn Ver. 2.0

<210> 1

<211> 468

<212> DNA

<213> primate

<220>

<221> misc_feature

<222> (301)

<223> nucleotide may be A, C, G, or T

<220>

<221> CDS

<222> (20)..(466)

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<221> mat_peptide

<222> (119)..(466)

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agtgtgaaac tggggtgga atg ggg tgt cca cgt atg ttc cct ttt gcc tta 52

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-30

-25

cta tat gtt ctg tca gtt tct ttc agg aaa atc ttc atc tta caa ctt 100

Leu Tyr Val Leu Ser Val Ser Phe Arg Lys Ile Phe Ile Leu Gln Leu

-20

-15

-10

gta ggg ctg gtg tta act tac gac ttc act aac tgt gac ttt gag aag 148

Val Gly Leu Val Leu Thr Tyr Asp Phe Thr Asn Cys Asp Phe Glu Lys

-5

-1

1

5

10

att aaa gca gcc tat ctc agt act att tct aaa gac ctg att aca tat 196

Ile Lys Ala Ala Tyr Leu Ser Thr Ile Ser Lys Asp Leu Ile Thr Tyr

15

20

25

atg agt ggg acc aaa agt acc gag ttc aac aac acc gtc tct tgt agc 244

Met Ser Gly Thr Lys Ser Thr Glu Phe Asn Asn Thr Val Ser Cys Ser

30

35

40

aat cgg cca cat tgc ctt act gaa atc cag agc cta acc ttc aat ccc 292

Asn Arg Pro His Cys Leu Thr Glu Ile Gln Ser Leu Thr Phe Asn Pro

45

50

55

aac cgc cgn gtg cgg tgc ctc gcc aaa gaa atg ttc gcc atg aaa act 340
 Asn Arg Xaa Val Arg Ser Leu Ala Lys Glu Met Phe Ala Met Lys Thr
 60 65 70

aag gct gcc tta gct atc tgg tgc cca ggc tat tgc gaa act cag ata 388
 Lys Ala Ala Leu Ala Ile Trp Cys Pro Gly Tyr Ser Glu Thr Gln Ile
 75 80 85 90

aat gct act cag gca atg aag aag agg aga aaa agg aaa gtc aca acc 436
 Asn Ala Thr Gln Ala Met Lys Lys Arg Lys Arg Lys Val Thr Thr
 95 100 105

aat aaa tgt ctg gaa caa gtg tca caa tta aa 468
 Asn Lys Cys Leu Glu Gln Val Ser Gln Leu
 110 115

<210> 2
 <211> 149
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 <222> (61)
 <223> residue will depend on translation of genetic code

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Val Ser Phe Arg Lys Ile Phe Ile Leu Gln Leu Val Gly Leu Val Leu
 -15 -10 -5

Thr Tyr Asp Phe Thr Asn Cys Asp Phe Glu Lys Ile Lys Ala Ala Tyr
 -1 1 5 10 15

Leu Ser Thr Ile Ser Lys Asp Leu Ile Thr Tyr Met Ser Gly Thr Lys
 20 25 30

Ser Thr Glu Phe Asn Asn Thr Val Ser Cys Ser Asn Arg Pro His Cys
 35 40 45

Leu Thr Glu Ile Gln Ser Leu Thr Phe Asn Pro Asn Arg Xaa Val Arg
 50 55 60

Ser Leu Ala Lys Glu Met Phe Ala Met Lys Thr Lys Ala Ala Leu Ala
 65 70 75

Ile Trp Cys Pro Gly Tyr Ser Glu Thr Gln Ile Asn Ala Thr Gln Ala
 80 85 90 95

Met Lys Lys Arg Arg Lys Arg Lys Val Thr Thr Asn Lys Cys Leu Glu
 100 105 110

Gln Val Ser Gln Leu
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<210> 3
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<220>
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 <222> (1)..(477)

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 atc ttc atc tta caa ctt gta ggg ctg gtg tta act tac gac ttc act 96
 Ile Phe Ile Leu Gln Leu Val Gly Leu Val Leu Thr Tyr Asp Phe Thr
 -10 -5 -1 1
 aac tgt gac ttt gag aag att aaa gca gcc tat ctc agt act att tct 144
 Asn Cys Asp Phe Glu Lys Ile Lys Ala Ala Tyr Leu Ser Thr Ile Ser
 5 10 15 20
 aaa gac ctg att aca tat atg agt ggg acc aaa agt acc gag ttc aac 192
 Lys Asp Leu Ile Thr Tyr Met Ser Gly Thr Lys Ser Thr Glu Phe Asn
 25 30 35
 aac acc gtc tct tgt agc aat cgg cca cat tgc ctt act gaa atc cag 240
 Asn Thr Val Ser Cys Ser Asn Arg Pro His Cys Leu Thr Glu Ile Gln
 40 45 50
 agc cta acc ttc aat ccc acc gcc ggc tgc gcg tcg ctc gcc aaa gaa 288
 Ser Leu Thr Phe Asn Pro Thr Ala Gly Cys Ala Ser Leu Ala Lys Glu
 55 60 65
 atg ttc gcc atg aaa act aag gct gcc tta gct atc tgg tgc cca ggc 336
 Met Phe Ala Met Lys Thr Lys Ala Ala Leu Ala Ile Trp Cys Pro Gly
 70 75 80
 tat tcg gaa act cag ata aat gct act cag gca atg aag aag agg aga 384
 Tyr Ser Glu Thr Gln Ile Asn Ala Thr Gln Ala Met Lys Lys Arg Arg
 85 90 95 100
 aaa agg aaa gtc aca acc aat aaa tgt ctg gaa caa gtg tca caa tta 432
 Lys Arg Lys Val Thr Thr Asn Lys Cys Leu Glu Gln Val Ser Gln Leu
 105 110 115
 caa gga ttg tgg cgt cgc ttc aat cga cct tta ctg aaa caa cag taa 480
 Gln Gly Leu Trp Arg Arg Phe Asn Arg Pro Leu Leu Lys Gln Gln
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Asn Cys Asp Phe Glu Lys Ile Lys Ala Ala Tyr Leu Ser Thr Ile Ser
5 10 15 20

Lys Asp Leu Ile Thr Tyr Met Ser Gly Thr Lys Ser Thr Glu Phe Asn
25 30 35

Asn Thr Val Ser Cys Ser Asn Arg Pro His Cys Leu Thr Glu Ile Gln
40 45 50

Ser Leu Thr Phe Asn Pro Thr Ala Gly Cys Ala Ser Leu Ala Lys Glu
55 60 65

Met Phe Ala Met Lys Thr Lys Ala Ala Leu Ala Ile Trp Cys Pro Gly
70 75 80

Tyr Ser Glu Thr Gln Ile Asn Ala Thr Gln Ala Met Lys Lys Arg Arg
85 90 95 100

Lys Arg Lys Val Thr Thr Asn Lys Cys Leu Glu Gln Val Ser Gln Leu
105 110 115

Gln Gly Leu Trp Arg Arg Phe Asn Arg Pro Leu Leu Lys Gln Gln
120 125 130

<210> 5

<211> 176

<212> PRT

<213> artiodactyla

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Asp Gly Gly Ala Tyr Gln Asn Val Leu Met Val Ser Ile Asp Asp Leu
35 40 45

Asp Asn Met Ile Asn Phe Asp Ser Asn Cys Leu Asn Asn Glu Pro Asn
50 55 60

Phe Phe Lys Lys His Ser Cys Asp Asp Asn Lys Glu Ala Ser Phe Leu
65 70 75 80

Asn Arg Ala Ala Arg Lys Leu Lys Gln Phe Leu Lys Met Asn Ile Ser
85 90 95

Asp Asp Phe Lys Leu His Leu Ser Thr Val Ser Gln Gly Thr Leu Thr
 100 105 110
 Leu Leu Asn Cys Thr Ser Lys Gly Lys Gly Arg Lys Pro Pro Ser Leu
 115 120 125
 Gly Glu Ala Gln Pro Thr Lys Asn Leu Glu Glu Asn Lys Ser Leu Lys
 130 135 140
 Glu Gln Arg Lys Gln Asn Asp Leu Cys Phe Leu Lys Ile Leu Leu Gln
 145 150 155 160
 Lys Ile Lys Thr Cys Trp Asn Lys Ile Leu Arg Gly Ile Thr Glu His
 165 170 175

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 Asp Gly Gly Ala Tyr Gln Asn Val Leu Met Val Asn Ile Asp Asp Leu
 35 40 45
 Asp Asn Met Ile Asn Phe Asp Ser Asn Cys Leu Asn Asn Glu Pro Asn
 50 55 60
 Phe Phe Lys Lys His Ser Cys Asp Asp Asn Lys Glu Ala Ser Phe Leu
 65 70 75 80
 Asn Arg Ala Ser Arg Lys Leu Arg Gln Phe Leu Lys Met Asn Ile Ser
 85 90 95
 Asp Asp Phe Lys Leu His Leu Ser Thr Val Ser Gln Gly Thr Leu Thr
 100 105 110
 Leu Leu Asn Cys Thr Ser Lys Gly Lys Gly Arg Lys Pro Pro Ser Leu
 115 120 125
 Ser Glu Ala Gln Pro Thr Lys Asn Leu Glu Glu Asn Lys Ser Ser Lys
 130 135 140
 Glu Gln Lys Lys Gln Asn Asp Leu Cys Phe Leu Lys Ile Leu Leu Gln
 145 150 155 160
 Lys Ile Lys Thr Cys Trp Asn Lys Ile Leu Arg Gly Ile Lys Glu His
 165 170 175

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<212> PRT
<213> primate

<400> 7
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Asp Gly Lys Gln Tyr Glu Ser Val Leu Met Val Ser Ile Asp Gln Leu
35 40 45
Leu Asp Ser Met Lys Glu Ile Gly Ser Asn Cys Leu Asn Asn Glu Phe
50 55 60
Asn Phe Phe Lys Arg His Ile Cys Asp Ala Asn Lys Glu Gly Met Phe
65 70 75 80
Leu Phe Arg Ala Ala Arg Lys Leu Arg Gln Phe Leu Lys Met Asn Ser
85 90 95
Thr Gly Asp Phe Asp Leu His Leu Leu Lys Val Ser Glu Gly Thr Thr
100 105 110
Ile Leu Leu Asn Cys Thr Gly Gln Val Lys Gly Arg Lys Pro Ala Ala
115 120 125
Leu Gly Glu Ala Gln Pro Thr Lys Ser Leu Glu Glu Asn Lys Ser Leu
130 135 140
Lys Glu Gln Lys Lys Leu Asn Asp Leu Cys Phe Leu Lys Arg Leu Leu
145 150 155 160
Gln Glu Ile Lys Thr Cys Trp Asn Lys Ile Leu Met Gly Thr Lys Glu
165 170 175

His

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<212> PRT
<213> rodent

<400> 8
Met Phe His Val Ser Phe Arg Tyr Ile Phe Gly Ile Pro Pro Leu Ile
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Leu Val Leu Leu Pro Val Thr Ser Ser Glu Cys His Ile Lys Asp Lys
20 25 30

130

135

140

Cys Trp Asn Lys Ile Leu Lys Gly Ser Ile
145 150